

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/517,214A
Source: Per
Date Processed by STIC: 2/9/06

ENTERED



PC

RAW SEQUENCE LISTING

DATE: 02/09/2006

PATENT APPLICATION: US/10/517,214A

TIME: 12:11:33

Input Set : A:\3065 USOP seq V2.txt

Output Set: N:\CRF4\02012006\J517214A.raw

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3 <110> APPLICANT: MAEKAWA, Tsuyoshi
4     HARA, Ryoma
5     ODAKA, Hiroyuki
6     MIZUFUNE, Hideya
7     FUKATSU, Kohji
9 <120> TITLE OF INVENTION: 1,2-Azole Derivatives with Hypoglycemic and Hypolipidemic Activity
11 <130> FILE REFERENCE: 3065 USOP
13 <140> CURRENT APPLICATION NUMBER: US/10/517,214A
14 <141> CURRENT FILING DATE: 2004-11-23
16 <150> PRIOR APPLICATION NUMBER: PCT/JP03/06389
17 <151> PRIOR FILING DATE: 2003-05-22
19 <150> PRIOR APPLICATION NUMBER: JP 2002-151405
20 <151> PRIOR FILING DATE: 2002-05-24
22 <150> PRIOR APPLICATION NUMBER: JP 2002-287161
23 <151> PRIOR FILING DATE: 2002-09-30
25 <150> PRIOR APPLICATION NUMBER: JP 2003-016748
26 <151> PRIOR FILING DATE: 2003-01-24
28 <160> NUMBER OF SEQ ID NOS: 12
30 <210> SEQ ID NO: 1
31 <211> LENGTH: 34
32 <212> TYPE: DNA
34 <213> ORGANISM: Artificial
-> 35 <220> FEATURE:
36 <223> OTHER INFORMATION: Primer/probe
38 <400> SEQUENCE: 1
39 aacggtacct cagccatgga gcagcctcag gagg      34
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42 <211> LENGTH: 34
43 <212> TYPE: DNA
45 <213> ORGANISM: Artificial
-> 46 <220> FEATURE:
47 <223> OTHER INFORMATION: Primer/probe
49 <400> SEQUENCE: 2
50 taagtcgacc cgtagtagta tgtccttgta gatc      34
52 <210> SEQ ID NO: 3
53 <211> LENGTH: 33
54 <212> TYPE: DNA
56 <213> ORGANISM: Artificial
-> 57 <220> FEATURE:
58 <223> OTHER INFORMATION: Primer/probe
60 <400> SEQUENCE: 3
61 ttagaattcg acatggacac caaacatttc ctg      33
63 <210> SEQ ID NO: 4

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64 <211> LENGTH: 33
65 <212> TYPE: DNA
67 <213> ORGANISM: Artificial
-> 68 <220> FEATURE:
69 <223> OTHER INFORMATION: Primer/probe
71 <400> SEQUENCE: 4
72 cccctcgagc taagtcattt ggtgcggcgc ctc 33
74 <210> SEQ ID NO: 5
75 <211> LENGTH: 36
76 <212> TYPE: DNA
78 <213> ORGANISM: Artificial
-> 79 <220> FEATURE:
80 <223> OTHER INFORMATION: Primer/probe
82 <400> SEQUENCE: 5
83 tcgacagggg accaggacaa aggtcacgtt cgggag 36
85 <210> SEQ ID NO: 6
86 <211> LENGTH: 36
87 <212> TYPE: DNA
89 <213> ORGANISM: Artificial
-> 90 <220> FEATURE:
91 <223> OTHER INFORMATION: Primer/probe
93 <400> SEQUENCE: 6
94 tcgactcccg aacgtgacct ttgtcctggt cccctg 36
96 <210> SEQ ID NO: 7
97 <211> LENGTH: 28
98 <212> TYPE: DNA
100 <213> ORGANISM: Artificial
-> 101 <220> FEATURE:
102 <223> OTHER INFORMATION: Primer/probe
104 <400> SEQUENCE: 7
105 cccagatctc cccagcgtct tgtcattg 28
107 <210> SEQ ID NO: 8
108 <211> LENGTH: 28
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
-> 111 <220> FEATURE:
112 <223> OTHER INFORMATION: Primer/probe
114 <400> SEQUENCE: 8
115 tcaccatggt caagctttta agcggggtc 28
117 <210> SEQ ID NO: 9
118 <211> LENGTH: 33
119 <212> TYPE: DNA
121 <213> ORGANISM: Artificial
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125 <400> SEQUENCE: 9
126 gtgggtaccg aaatgaccat ggttgacaca gag 33
128 <210> SEQ ID NO: 10
129 <211> LENGTH: 33

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RAW SEQUENCE LISTING

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Input Set : A:\3065 US0P seq V2.txt

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130 <212> TYPE: DNA
132 <213> ORGANISM: Artificial
-> 133 <220> FEATURE:
134 <223> OTHER INFORMATION: Primer/probe
136 <400> SEQUENCE: 10
137 ggggtcgacc aggactctct gctagtacaa gtc 33
139 <210> SEQ ID NO: 11
140 <211> LENGTH: 33
141 <212> TYPE: DNA
143 <213> ORGANISM: Artificial
-> 144 <220> FEATURE:
145 <223> OTHER INFORMATION: Primer/probe
147 <400> SEQUENCE: 11
148 cgtcgacccg gcggcccat ggacctgccc ccg 33
150 <210> SEQ ID NO: 12
151 <211> LENGTH: 33
152 <212> TYPE: DNA
154 <213> ORGANISM: Artificial
-> 155 <220> FEATURE:
156 <223> OTHER INFORMATION: Primer/probe
158 <400> SEQUENCE: 12
159 catcgattag cagtggcggtt acttctggga ctt 33

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/517,214A

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valid <213> Response:

is of "Artificial" only as "<213> Organism" response is incomplete,
r 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

seq#:1,2,3,4,5,6,7,9,10,11,12

VERIFICATION SUMMARY

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Input Set : A:\3065 USOP seq V2.txt

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35 M:283 W: Missing Blank Line separator, <220> field identifier
46 M:283 W: Missing Blank Line separator, <220> field identifier
57 M:283 W: Missing Blank Line separator, <220> field identifier
58 M:283 W: Missing Blank Line separator, <220> field identifier
79 M:283 W: Missing Blank Line separator, <220> field identifier
90 M:283 W: Missing Blank Line separator, <220> field identifier
101 M:283 W: Missing Blank Line separator, <220> field identifier
111 M:283 W: Missing Blank Line separator, <220> field identifier
122 M:283 W: Missing Blank Line separator, <220> field identifier
133 M:283 W: Missing Blank Line separator, <220> field identifier
144 M:283 W: Missing Blank Line separator, <220> field identifier
155 M:283 W: Missing Blank Line separator, <220> field identifier